

FIG. 1

CCCAGTGCCA TTTTCTCTCT CTAGTCAAGC TCTCTATATC ATCATCACTA TTCCCTTGGC
 TGCAGTAACA CTCCTATTTA ACCCTCACAA AAAAATTACC AGAGGGCAGC AAAAAATGCT
 TGAACATAAT TATTATACTT ACTATTAAGC TAGATTTCCCT CTTGATCTTG CTAGGTTTGA
 CTGGAGAAAA TGGCAGGCAT GGATAGAAAC AGTTTCAACA GTAAGTACTT CAAAAACAAA
 M A G M D R N S F N S K Y F K N K
 AGCATCATGG CAAGACAGAT GGAGTACTTG AATAACAACA ATGGCGACAA TAACAACAAC
 S I M A R Q M E Y L N N N N G D N N N N
 AATAATGTTA CAAGCTCATT ACGAGATAAT TATGGAAATG AAGATCATT ACTTGGTGGG
 N N V T S S L R D N Y G N E D H L L G G
 CTATTCTCTT GGCCTCCAAG ATCTTATACA TGTAGCTTTT GTAAAAGGGA ATTTAGATCT
 L F S W P P R S Y T C S F C K R E F R S
 GCTCAAGCTC TTGGTGGACA CATGAATGTT CATAGAAGAG ATAGAGCCAT TTTGAGACAA
 A Q A L G G H M N V H R R D R A I L R Q
 TCACCACCTA GAGATATTAA TAGGTATTCT CTTCTAAACC TTAATCTTGA ACCAAACCCT
 S P P R D I N R Y S L L N L N L E P N P
 AACTTTTACC CTAGTCATAA CCCTAGTTTT TCAAGAAAAT TCCCACCTTT TGAAATGAGG
 N F Y P S H N P S F S R K F P P F E M R
 AAATTAGGAA AAGGAGTTGT TCCAAACAAT CACTTGAAAA GTGCCAGAGG GCGTTTTGGA
 K L G K G V V P N N H L K S A R G R F G
 GTTGAGAAAA TTGACTCTTT CATGCAAGAA AAAGAATGTA CTACTACAGT GATCAAGAAG
 V E K I D S F M Q E K E C T T T V I K K
 TCCGAGTTTC TAAGATTGGA CTTGGGAATT GGGTTGATCA GTGAATCAAA GGAAGATTTA
 S E F L R L D L G I G L I S E S K E D L
 GATCTTGAAC TTCGACTGGG ATCCACTTAA CTATATCTAA TTTTACGGC ATTAAGGTTT
 D L E L R L G S T
 GTAAATTGAG TCGACAGCTT AGTCAAACT ACTTATGCAC TTTAATATGG CTTCTTGTGC
 TATATTTATT TATTTTACAT GGCTGTATCT AGGTTTGCAT TTTAAGATTT AGTACCTTGT
 CAGATTAAAA GAAAACGAAA GTTAAATTAA AAAAAA

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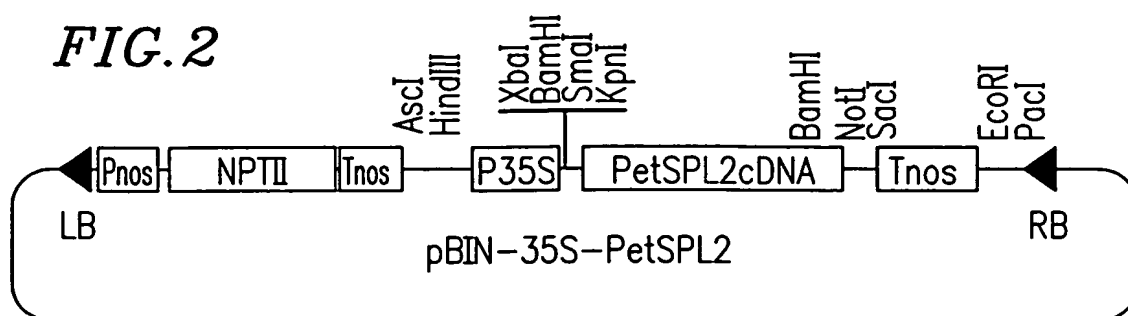
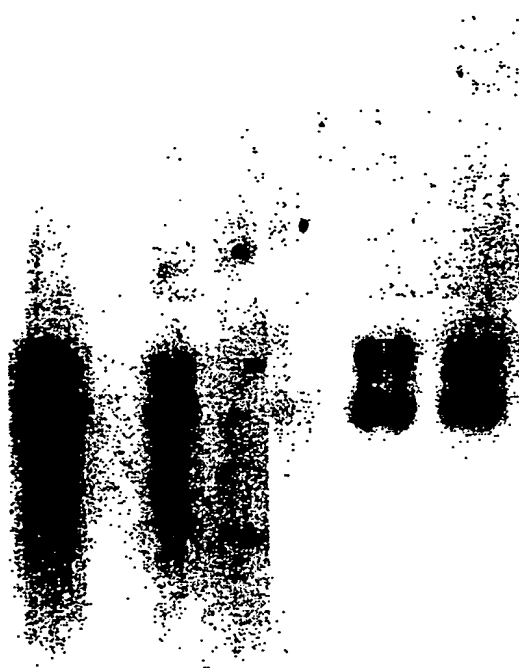
FIG. 2

FIG. 3

1 2 3 4 5 6 7 8 9 10 11 12 13



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FIG. 4



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FIG. 5



868T60" 08595T60